

ERRATUM: Payseur BA, Covert HA, Vinyard CJ, Dagosto M. 1999. New Body Mass Estimates for *Omomys carteri*, a Middle Eocene Primate From North America. Am J Phys Anthropol 109:41–52.

This article included an incomplete Table 2. The final two columns, showing “Intercept” and “SEE” data were omitted. The complete Table 2, with these two columns included, is provided below.

TABLE 2. *O. carteri* body mass estimates and associated statistics from bivariate regression analyses¹

Character	<i>n</i>	Dataset	Body mass estimate (g)	Upper CL	Lower CL	<i>R</i> ²	Slope	Intercept	SEE
SGH	1	AP	111.7	132.1	94.5	0.951	2.716	1.044	0.304
		S	113.0	150.8	84.7	0.917	2.726	1.023	0.362
		H	111.2	138.8	89.2	0.964	2.708	1.062	0.263
SGW	1	AP	251.9	286.5	221.5	0.948	2.503	2.731	0.311
		S	266.6	331.2	214.6	0.905	2.627	2.624	0.389
		H	230.6	267.8	198.6	0.973	2.514	2.653	0.227
HHW	3	AP	252.1	289.7	219.3	0.940	2.646	1.323	0.335
		S	269.6	348.0	208.9	0.872	2.751	1.180	0.451
		H	230.6	263.8	201.6	0.978	2.655	1.255	0.205
HHH	5	AP	277.3	316.6	242.9	0.942	2.852	0.814	0.331
		S	273.7	341.0	219.8	0.901	2.900	0.697	0.397
		H	285.4	339.1	240.2	0.959	2.803	0.940	0.280
HDA	1	AP	183.9	206.3	163.9	0.966	2.703	2.600	0.251
		S	175.8	215.1	143.6	0.940	2.693	2.654	0.309
		H	169.2	195.6	146.4	0.980	2.753	2.481	0.199
HDM	2	AP	356.9	393.3	323.9	0.962	2.589	2.673	0.269
		S	250.9	283.3	222.2	0.934	2.661	2.237	0.251
		H	345.0	392.2	303.5	0.974	2.593	2.645	0.225
HAW	6	AP	408.6	449.3	371.6	0.959	2.616	0.987	0.276
		S	373.0	418.5	332.5	0.963	2.670	0.802	0.241
		H	451.5	526.1	387.5	0.956	2.530	1.246	0.293
HBW	2	AP	334.7	374.1	299.4	0.952	2.781	−0.542	0.300
		S	320.3	376.0	272.8	0.938	2.749	−0.518	0.315
		H	122.8	144.9	104.0	0.956	2.282	−0.410	0.291
RHM	2	AP	255.5	282.2	231.4	0.968	2.645	2.200	0.244
		S	274.2	322.4	233.2	0.943	2.646	2.254	0.300
		H	232.2	262.3	205.5	0.982	2.706	2.040	0.187
RDM	1	AP	262.6	295.5	233.3	0.955	2.599	1.861	0.291
		S	285.0	344.6	235.7	0.921	2.560	1.978	0.353
		H	235.5	274.3	202.2	0.972	2.680	1.653	0.233
USNW	2	AP	281.3	319.9	247.2	0.944	2.624	2.576	0.324
		S	277.6	340.0	226.7	0.913	2.608	2.565	0.371
		H	290.6	347.4	243.1	0.956	2.603	2.643	0.291
USNH ²	1	AP	157.4	179.3	138.2	0.962	2.449	2.292	0.268
		S	166.1	203.6	135.4	0.941	2.517	2.259	0.305
		H	140.5	167.1	118.1	0.974	2.494	2.139	0.223
PAH ²	2	AP	156.0	183.2	132.7	0.943	2.700	0.651	0.326
		S	137.5	175.3	107.9	0.930	2.654	0.597	0.333
		H	183.1	221.8	151.2	0.963	2.631	0.940	0.267
PAW	1	AP	202.6	229.6	178.7	0.958	2.577	1.124	0.281
		S	137.5	169.8	111.4	0.937	2.654	0.597	0.333
		H	218.7	259.4	184.5	0.967	2.502	1.329	0.253
PILL ³	1	AP	221.5	252.0	194.7	0.953	2.618	−2.920	0.298
		S	173.6	204.8	147.1	0.959	2.910	−4.075	0.255
		H	271.6	322.5	228.8	0.961	2.424	−2.096	0.274
PISL ²	1	AP	389.2	457.7	331.0	0.894	2.318	0.153	0.447
		S	425.2	477.0	379.0	0.961	2.581	−0.338	0.250
		H	311.2	425.4	227.6	0.874	2.454	−0.428	0.494

(Continued)

TABLE 2. (continued)

Character	<i>n</i>	Dataset	Body mass estimate (g)	Upper CL	Lower CL	R^2	Slope	Intercept	SEE
FHH	8	AP	244.3	271.9	219.4	0.964	2.676	1.500	0.259
		S	247.2	289.1	211.3	0.951	2.548	1.696	0.279
FHA	5	H	242.7	283.4	207.9	0.971	2.729	1.420	0.239
		AP	257.9	297.4	223.7	0.937	2.695	1.488	0.345
FCW	4	S	274.6	340.1	221.8	0.905	2.405	1.966	0.388
		H	235.0	280.7	196.7	0.962	2.886	1.134	0.270
FCH ²	4	AP	304.0	333.6	277.0	0.968	2.653	0.105	0.244
		S	303.9	348.1	265.4	0.956	2.816	-0.243	0.263
FDA	2	H	302.2	345.3	264.5	0.974	2.604	0.207	0.223
		AP	673.6	771.8	587.9	0.893	2.876	-0.093	0.449
FDM	2	S	524.4	628.4	437.6	0.896	2.622	0.249	0.407
		H	784.7	920.8	668.6	0.931	2.944	-0.060	0.365
TPW	3	AP	331.2	369.4	296.8	0.954	2.671	2.338	0.294
		S	292.1	331.4	257.4	0.918	2.661	2.237	0.251
TPA	3	H	324.6	370.8	284.2	0.973	2.736	2.252	0.228
		AP	230.2	254.3	208.4	0.970	2.724	2.159	0.236
TDA ²	3	S	271.4	333.7	220.7	0.960	2.526	2.523	0.360
		H	229.9	267.4	197.7	0.973	2.746	2.133	0.229
PCA-Corr.	—	AP	339.1	370.6	310.2	0.968	2.664	0.049	0.243
		S	305.8	346.5	269.9	0.963	2.794	-0.335	0.244
PCA-Cov.	—	H	378.7	428.2	334.9	0.974	2.544	0.423	0.223
		AP	539.9	611.6	476.6	0.919	2.470	1.611	0.390
PCA-Cov.	—	S	555.9	694.4	445.0	0.842	2.280	1.945	0.501
		H	514.6	589.6	449.1	0.961	2.569	1.417	0.274
PCA-Cov.	—	AP	252.9	295.9	216.1	0.925	2.757	1.746	0.375
		S	186.8	229.0	152.4	0.936	2.862	1.321	0.319
PCA-Cov.	—	H	325.5	387.3	273.6	0.955	2.600	2.237	0.295
		AP	252.0	275.4	230.6	0.975	1.340	6.888	0.217
PCA-Cov.	—	S	247.7	283.6	216.3	0.962	1.207	6.437	0.245
		H	276.5	312.1	245.0	0.980	1.351	7.246	0.199
PCA-Cov.	—	AP	329.7	286.8	379.1	0.975	0.543	6.888	0.215
		S	245.0	213.9	280.5	0.961	0.549	6.437	0.242
PCA-Cov.	—	H	393.7	323.2	479.6	0.979	0.537	7.246	0.198

¹ *n* = number of *O. carteri* specimens from which the species mean was estimated; AP = all-primates; S = strepsirrhines; H = haplorhines; Upper CL = 95% upper confidence limit; Lower CL = 95% lower confidence limit; R^2 = coefficient of determination; SEE = standard error of the estimate; PCA-Corr = species scores for first component in a principal component analysis on the correlation matrix of all variables; PCA-Cov = species scores for the first component in a principal components analysis on the variance-covariance matrix of all variables; Regression statistics are the results of analyses using *ln*-transformed values.

² Strepsirrhine and haplorhine slopes are significantly different ($P < 0.05$).

³ Strepsirrhine and haplorhine intercepts are significantly different ($P < 0.05$).